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				つりに					აას				Phe	000		
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	0.77	Are	g Arg			- 375					300	,			Gly	
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tca Sei	a ago r Sei	c tto c Phe	c tta e Lei 20	ı Thi	tca Sei	a aac Asi	acc n Thi	c ctt r Lei 25	і Суя	tca Se:	a gat r Asj	gaa Gli	a tta ı Len 30	ı Med	atg : Met	96
cc Pr	c ca	t tt s Pho	e Hi	s Se	r Lys	5 GL	ı Gı	t tat y Tyi	c GI	a aa y Ly	a ta s Ty:	t ta r Ty: 4	. 61	g ctg n Lei	g aga ı Arg	144
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Thr Ala Arg Arg Ile Thr Lys Thr Leu Ala Gly Leu Pro Gln Lys Ser
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Phe Thr Glu Thr Asp Asp Ala Glu Arg Lys Gln Glu Lys Ile Gly Asn
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cca Pro	gcc Ala	cct Pro	gcc Ala 80	aac Asn	aaa Lys	gtg Val	ccc Pro	cac His 85	tca Ser	gca Ala	gcc Ala	aac Asn	ctg Leu 90	ccc Pro	ctg Leu	292
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cag Gln	aaa Lys	ccc Pro	ctg Leu	cac His 145	tca Ser	ctg Leu	ggc	Ser	agc Ser 150	gag Glu	ttg Leu	ctc Leu	tac Tyr	gtc Val 155	atg Met	484
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Ala Tyr Val Leu Ile Phe Leu Leu Cys Met Val Gly Asn Thr Leu Val 50 55 60

Cys Phe Ile Val Leu Lys Asn Arg His Met Arg Thr Val Thr Asn Met 65 70 75 80

Phe Ile Leu Asn Leu Ala Val Ser Asp Leu Leu Val Gly Ile Phe Cys 85 90 95

Met Pro Thr Thr Leu Val Asp Asn Leu Ile Thr Gly Trp Pro Phe Asp 100 105 110

Asn Ala Thr Cys Lys Met Ser Gly Leu Val Gln Gly Met Ser Val Ser 115 120 125

Ala Ser Val Phe Thr Leu Val Ala Ile Ala Val Glu Arg Phe Arg Cys

Ile Val His Pro Phe Arg Glu Lys Leu Thr Leu Arg Lys Ala Leu Phe
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Thr Ile Ala Val Ile Trp Ala Leu Ala Leu Leu Ile Met Cys Pro Ser

Ala Val Thr Leu Thr Val Thr Arg Glu Glu His His Phe Met Leu Asp 180 185 190

Ala Arg Asn Arg Ser Tyr Pro Leu Tyr Ser Cys Trp Glu Ala Trp Pro

Glu Lys Gly Met Arg Lys Val Tyr Thr Ala Val Leu Phe Ala His Ile

Tyr Leu Val Pro Leu Ala Leu Ile Val Val Met Tyr Val Arg Ile Ala 225 230 235 240

Arg Lys Leu Cys Gln Ala Pro Gly Pro Ala Arg Asp Thr Glu Glu Ala

Val Ala Glu Gly Gly Arg Thr Ser Arg Arg Arg Ala Arg Val Val His 260 265 270 Met Leu Val Met Val Ala Leu Phe Phe Thr Leu Ser Trp Leu Pro Leu 280

Trp Val Leu Leu Leu Ile Asp Tyr Gly Glu Leu Ser Glu Leu Gln 295 290

Leu His Leu Leu Ser Val Tyr Ala Phe Pro Leu Ala His Trp Leu Ala 315 310

Phe Phe His Ser Ser Ala Asn Pro Ile Ile Tyr Gly Tyr Phe Asn Glu 330 325

Asn Phe Arg Arg Gly Phe Gln Ala Ala Phe Arg Ala Gln Leu Cys Trp 345

Pro Pro Trp Ala Ala His Lys Gln Ala Tyr Ser Glu Arg Pro Asn Arg 360 355

Leu Leu Arg Arg Arg Val Val Val Asp Val Gln Pro Ser Asp Ser Gly 375

Leu Pro Ser Glu Ser Gly Pro Ser Ser Gly Val Pro Gly Pro Gly Arg 395 390

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His Gln Glu Ile Gln Ser Pro Gly Gln Glu Gln Pro Arg Lys Arg Val 170 Phe Thr Glu Thr Asp Asp Ala Glu Arg Lys Gln Glu Lys Ile Gly Asn 185 Leu Gln Pro Val Leu Gln Gly Ala Met Lys Leu 200 <210> 51 <211> 609 <212> DNA <213> Rattus sp. <400> 51 atggaaatta tttcatcaaa gcgattcatt ttattgactt tagcaacttc aagcttctta 60 acttcaaaca ccctttgttc agatgaatta atgatgcccc attttcacag caaagaaggt 120 tatggaaaat attaccagct gagaggaatc ccaaaagggg taaaggaaag aagtgtcact 180 tttcaagaac tcaaagattg gggggcaaag aaagatatta agatgagtcc agcccctgcc 240 aacaaagtgc cccactcagc agccaacctt cccctgaggt ttgggaggaa catagaagac 300 agaagaagcc ccagggcacg ggccaacatg gaggcaggga ccatgagcca ttttcccagc 360 ctgccccaaa ggtttgggag aacaacagcc agacgcatca ccaagacact ggctggtttg 420 cccagaaat ccctgcactc cctggcctcc agtgaattgc tctatgccat gacccgccag 480 catcaagaaa ttcagagtcc tggtcaagag caacctagga aacgggtgtt cacggaaaca 540 gatgatgcag aaaggaaaca agaaaaata ggaaacctcc agccagtcct tcaaggggct 600 atgaagctg <210> 52 <211> 27 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Primer <400> 52 27 ttctagattt tggacaaaat ggaaatt <210> 53 <211> 27 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Primer <400> 53 27 cgtctttagg gacaggctcc agatttc

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295

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Ser 385	Glu	Ser	Gly	Pro	Ser 390	Ser	Gly	Ala	Pro	Arg 395	Pro	Gly	Arg	Leu	Pro 400
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